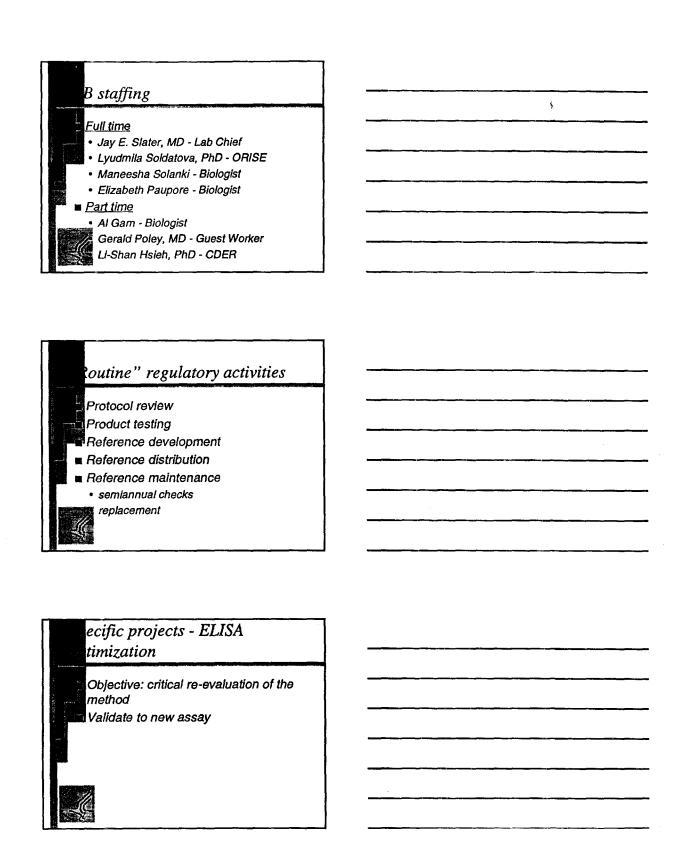
	ł
boratory of	
munobiochemistry	
Operational issues	
B - missions	
Research	
Product quality Regulatory support	
■ Industry support	
ting chiefs of LIB, 1997-98	
Paul Turkeltaub, MD	
Richard Pastor, PhD	



JSA optimization

Buffer detergent has been changed from Brij to Tween-20.

The blocking buffer and test diluent have been changed to PBS pH 7.4 containing 0.05% Tween-20 and 1.0 % bovine serum albumin

■ Coating, competition and conjugate incubations are all overnight.

MB substrate is equilibrated at room imperature for 5 min; TMB incubation step is actly 5 min.

ISA validation results

No 1.0 extracts failed No 0.5/2.0

No 0.5/2.0 extracts passed

SD within old limits (0.1375)

■ Accurate with all three antigens, at full range of concentrations

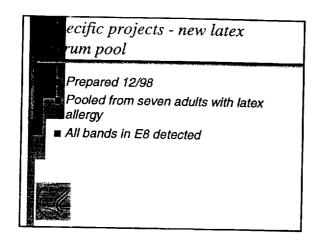
		N	m)	sd[log ip]
m eadow	0.5	24	0.516	0.097
fescue	1	24	1.104	0.107
	2	24	2.085	0.152
D. fadnae	0.5	23	0.499	0.109
	1	23	1.067	0.105
	2	23	2.19	0.113
Bermuda	0.5	23	0.464	0.116
	1	23	0.9914	0.102
	2	23	1.94	0.125

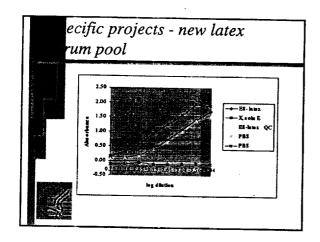
w reference sera/extracts

Cat S2 replaced by S2a Mite S3 replaced by S4 Latex S2 replaced by S3

■ Dp and cat extract replacements in progress

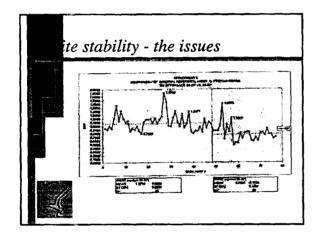


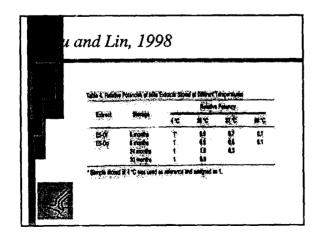




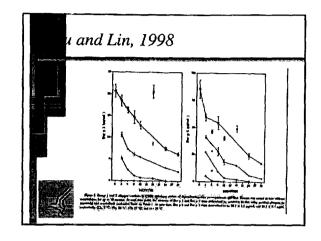
			in this to the contraction of	-o-uto-ina-ina-ina-ina-ina-ina-ina-ina-ina-ina	
		X, 1	mg/mL	E8, 3.9 m	g/mL
	Mean log RP		-0.63		0.04
34-45	SD (log RP)		0.05		0.05
	Mean RP		0.24	***************************************	1.10
İ	95% CI	0.19	to 0.30	0.89 to	

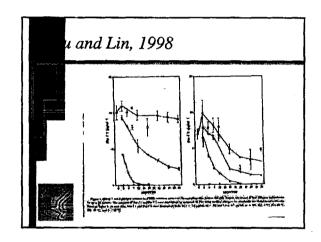
Cysteine and serine protease activity of mite antigens Conflicting prior data on stability Nelson et al. J Allergy Clin Immunol 1996;98:382. Liu and Lin. Ann Allergy Asthma Immunol 1998;80:177. Problems associated with short shelf life





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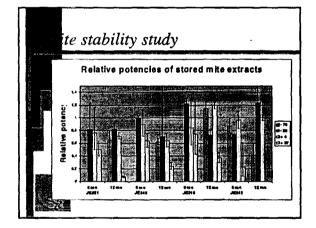
Der p	rther questi 1, Derp 2 and 2 unstable at	ONS Can protease inhibitors slow the
4°C ■ Der f	1 stable at	degradation? ■ Is RP conserved in
■ RP co	Instable >26°C Inserved at relative to 4°C	the 4°C specimen relative to lyophilized?

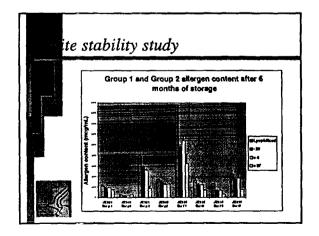
ite stability - <mark>experimental</mark> sign

Objective: identify and characterize possible degradation in glycerinated mite extracts, with and without protease inhibitors

Glycerinated extracts stored at -70°C, -20°C, +4°C and +37°C for 6-12 months

- Compare to lyophilized standard
- Assayed by three methods
 - · competitive ELIŞA for relative potency
 - two-site ELISA for group 1 and group 2 antigens
 Western blot (sera and monoclonals)

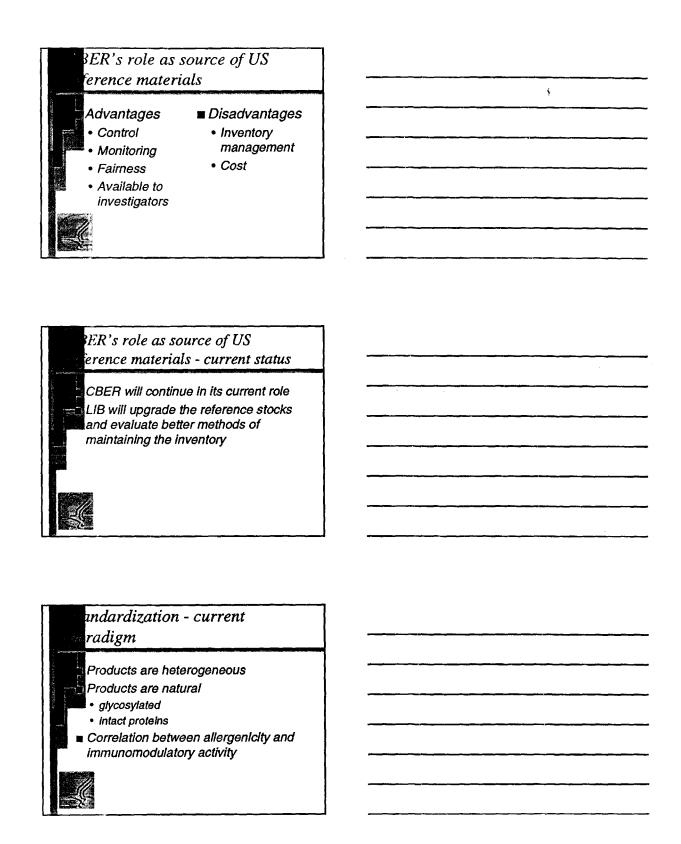




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, , , ,		

te stability study - tentative nclusions RP stable at 4°C relative to lyophilized loss of protein bands at 4°C loss of specific mite allergens at 4°C; this does not correlate with RP ■ protease inhibitors offer no long-term protection ference replacement program Current references: • many out of date (20/24) Replacement program: · bring full inventory up to date • target completion date: August 2001 ference replacement program Proactive - candidates will be identified >6 months prior to expiration Comprehensive - all reference materials will be updated ■ Anticipated problems - frequent replacement required possible solutions : lyophilized references; or ELISA based on serum pools

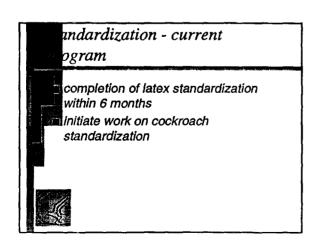
ference replacement program oposal 1998-99 : CBER will lyophilize a portion of all reference extracts/sera 1999-00 : CBER will assess stability and reliability of lyophilized products ■ Results and samples will be distributed to APMA membership prior to action ues for long term nsideration Should CBER continue to be the source of reference standard allergens and antisera? ■ How should the standardization program proceed? BER's role as source of US placement reference materials LIB identifies candidate reference • in house testing • samples sent to manufacturers for testing ■ LIB purchases 1-3 year supply ■ LIB distributes to manufacturers



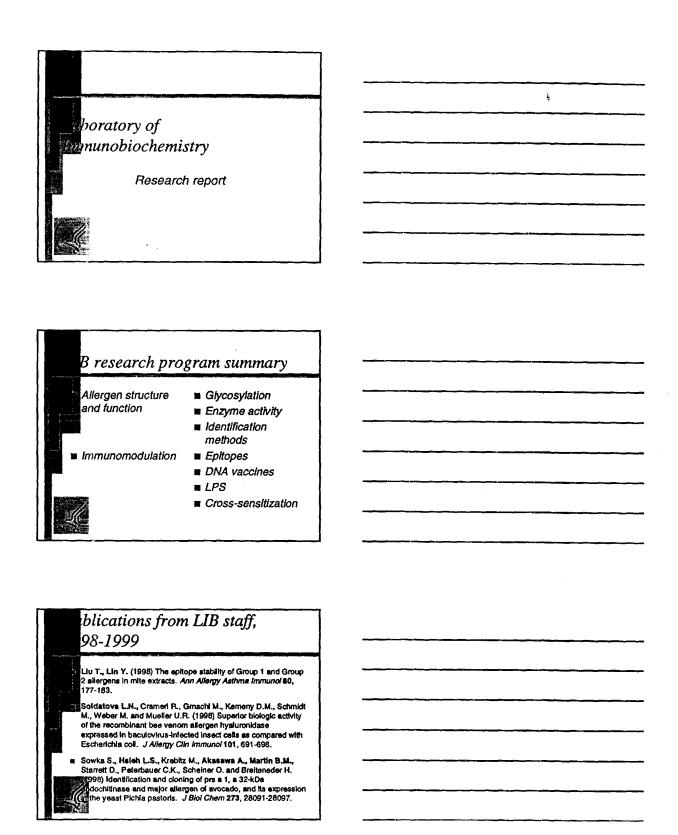
rrent standardization targets Latex Cockroach Tree pollens evea latex antigens Hev b 1 (REF) ■ hevamine Hev b 2 (β-1,3-■ chitinases (I and II) glucanase) ■ Mn-superoxide Hev b 3 (microhelix dismutase component) ■ enolase ■ Hev b 4 m profilin ■ Hev b 5 ■ lysozyme ■ Hev b 6 (prohevein) ev b 7 (patatin palogue) ■ proteasome subunit andardization - limitations Uncertain predictive value for peptides plasmids modified allergens · non-glycosylated products ■ Cost borne by FDA

Product	Source	Size	Cost
Filgrastim	E. coli	175 αα	\$0.53/µg
Sargramostim	Yeast	127 αα	\$0.52/µg
Epoetin-α	Chinese	165 αα	\$0.012/U

andardization - proaches	- alternative
1. Consistency monitoring 2. Pure allergen basis (monoclonal antibody) 3. Other in vitro characterization	■ <u>Disadvantages</u> ■ No industry standard ■ All allergens not identified or characterized ■ Criteria not established

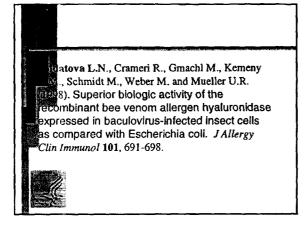


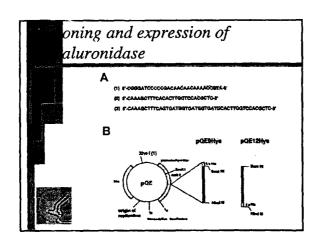
B staffing			
D Siujjing	ta), maaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa	-	<u> </u>
<u>Current</u>	■ <u>Proposed</u>		
Lab Chief	Add one more		
Post-doc	biologist]	
■ Two biologists			
■ One new biologist			
has been hired (to			
tart 3/1/99)			

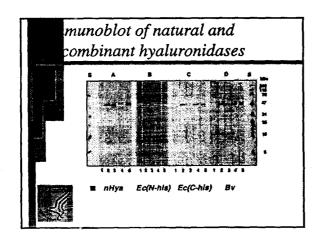


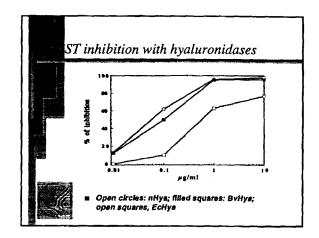
blications from LIB staff, 98-1999 Slater J.E., Paupore E., Zhang Y.T. and Colberg-Poley A.M. (1998) The latex allergen Hev b 5 transcript is widely distributed after subcutaneous injection in BALB/c mice of its DNA vaccine. J Allergy Clin Immunol 102, 469-475. Slater J.E., Paupore E.J., Elwell M.R. and Truscott W. (1998) Lipopolysaccharide augments igG and igE responses of mice to the latex allergen Hev b 5. J Allergy Clin Immunol 102, 977-Slater J.E., Paupore E.J. and O'Hehir R.E. (1999) Murine B-cell and T-cell epitopes of the allergen Hev b 5 from natural rubber latex. Molecular Immunology (In Press) lergen structure and function glycosylation Is the decreased antibody binding of nonglycosylated antigens primarily a function of Impaired folding? ■ What is the biochemical anatomy of the glycosylation requirement? ■ Can non-glycosylated allergens equal native allergens in immunotherapy? How can non-glycosylated products be valuated for diagnosis and therapy? lergen structure and function enzyme activity What is the relationship between enzyme activity and allergenicity? antibody binding - in vitro · bioavallability antigen processing ■ Specific regulatory applications

hymenoptera, mites, latex

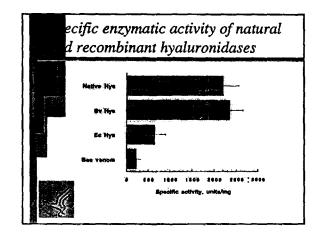




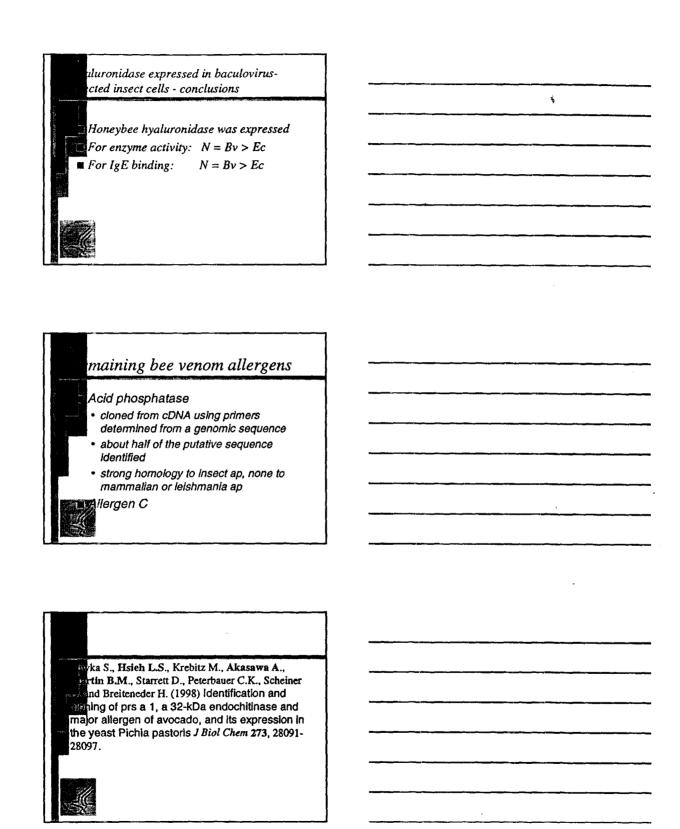


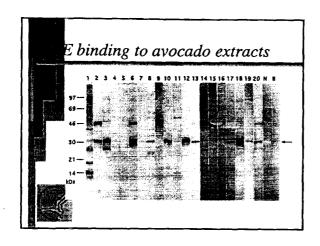


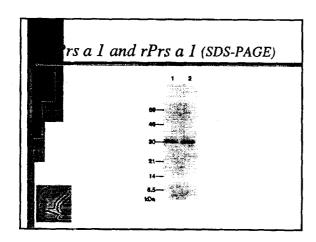
		ade a la	an american	
TABLE L Specific Scripp in 30 pulse norell engle dontro	te Merci	C to bee v	endm ew	30
	MART	- 40	offic left to	nu
Prihes! (4 - 90)	-	1(5)	1 (6)	1 (40)
	1 2 3	6 4 (24)	1 (15) 2 (15) 2 (15)	7(10) 7(15) 7(15)
	-	4 (28) 4 (28)	4 (45) 3 (18)	\$ (15)
Control subjects	3	36	20	-A-



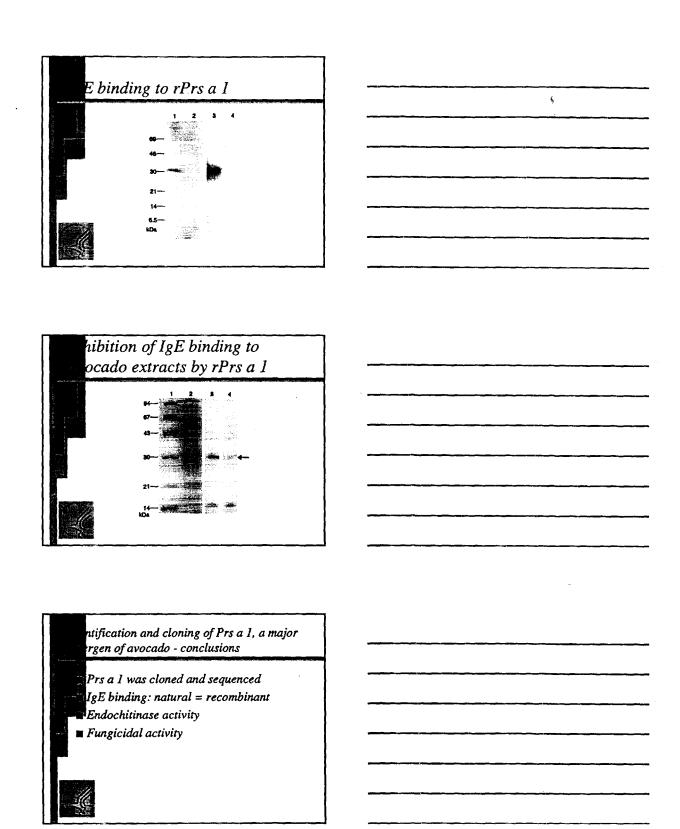
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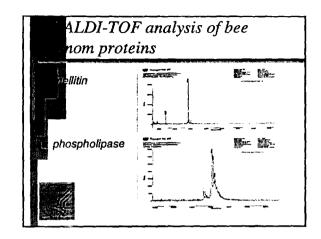


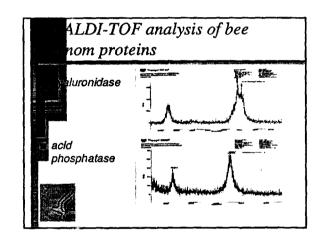


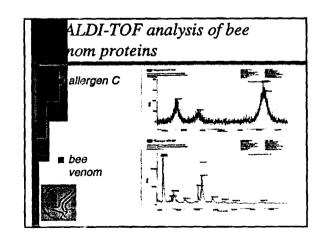
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to the first free course	A Pre a 1 1 BOCKROMARIATEREACUSQUENCESTEDUCCET-TODSCORE 41 [[[]]][[]][[]][[]][[]][[]][[]][[]][[]



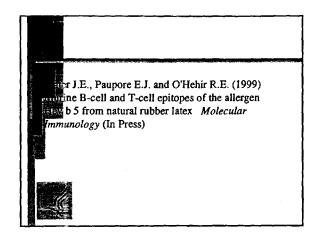
lergen structure and function ditional questions If an allergen that is not glycosylated, glycosylated abnormally, or denatured shows poor IgE binding or impaired enzymatic activity, how can we evaluate its efficacy as an immunotherapeutic reagent? lergen structure and function dentification methods MALDI-TOF Quantitative SDS-PAGE Quantitative immunoblot trix-<u>a</u>ssisted <u>l</u>aser <u>d</u>esorption/<u>i</u>onization -of-flight mass spectrometry (MALDI-TOF) Voyagar's Advanced Single Stage Reflector

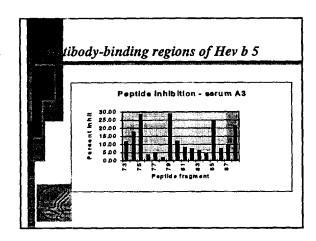


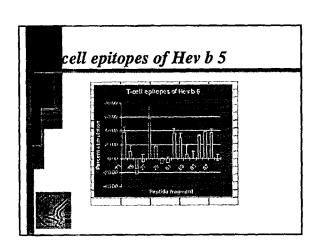


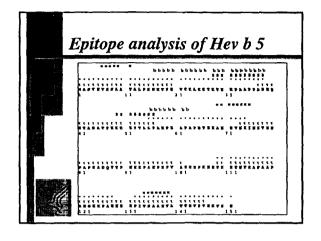


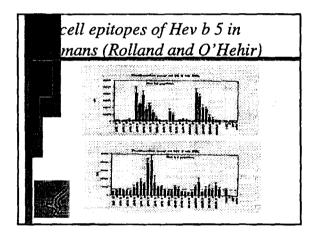
lergen identification techniques dditional questions	
Can we develop a quantitative profile of natural allergen preparations?	•
Can we use MALDI-TOF to carefully assess the glycosylation of recombinant	
allergens?	
munomodulation	
pitope specific therapy	
Human epitope analysis of Hev b 5 • Site directed mutagenesis	
Support for clinical trials of latex immunotherapeutic reagents	
	-
pitope-based immunotherapy	
To make a second and a second a	
Identify and purify antigen Identify T-cell epitopes of the	
antigen ■ Identify B-cell epitopes (IgE-	
binding sites) of the antigen	
Administer immunotherapy with he T-cell epitopes	
Tie 1-ceil epitopes	



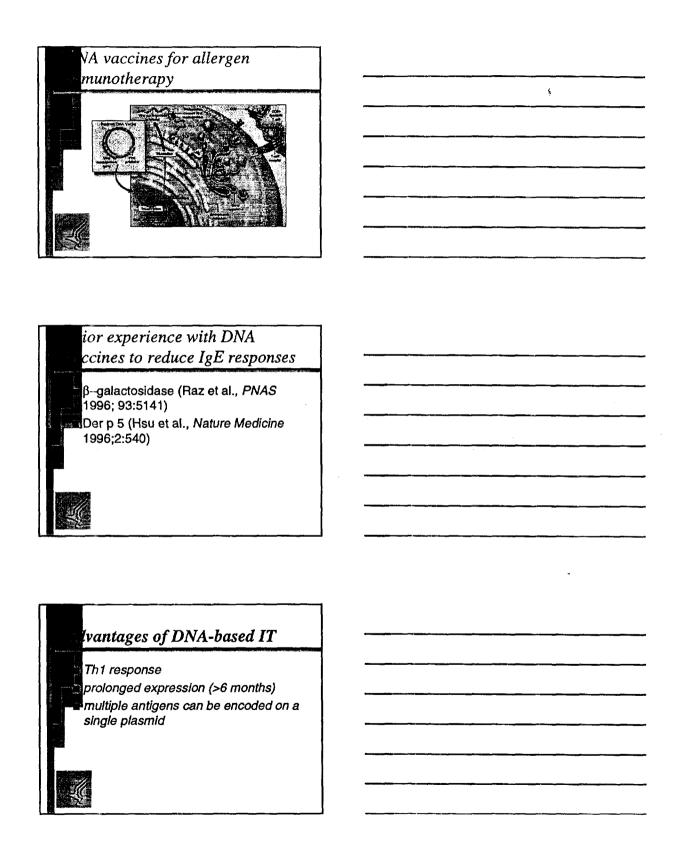




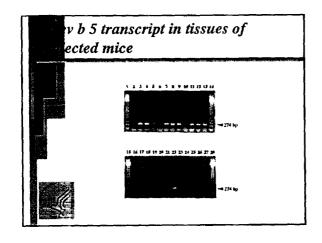


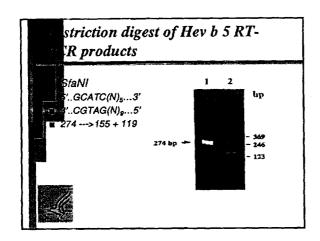


itopes of Hev b 5 - conclusions
Murine B-cell and T-cell epitopes of Hev b 5 have been identified
Preliminary identification of human T- cell epitopes suggests dominance (peptides 37-56, 73-92, 109-128 and 118-137)
Regions for mouse immunotherapy tudy identified

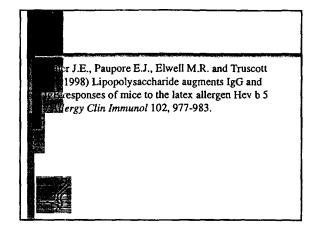


oblems with DNA-based IT unproven safety profile • mutagenesis · tissue specificity • allergen release • CD8 responses ■ control of responses in vivo eliminary experience with Hev DNA vaccine the sense construct is toxic to presensitized mice when injected into the tongue ■ no toxicity was noted when the construct was injected intradermally Slater J.E., Paupore E., Zhang Y.T. and Colberg-Poley A.M. (1998) The latex allergen Hev b 5 transcript is widely distributed after subcutaneous injection in BALB/c mice of its DNA vaccine J Allergy Clin Immunol 102, 469-475

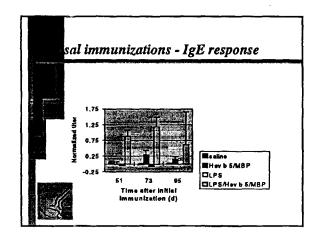


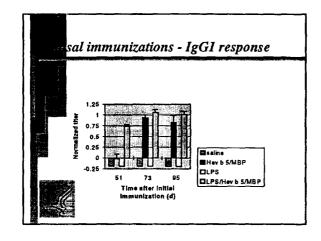


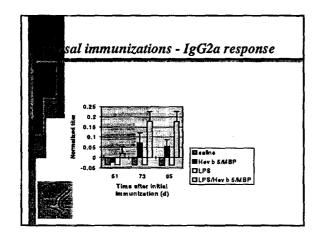
	IA vaccine therapy - further dies
The state of the s	Construction of DNA vaccines for Hev b 5 using • specific T-cell epitopes • weak promoters • tissue-specific promoters



sal immunization protocol BALB/c mice received either saline, LPS, Hev b 5, or LPS + Hev b 5 (10 μg each) Anesthetized with methoxyflurane Received qod doses on days 1-12 and days 64-68

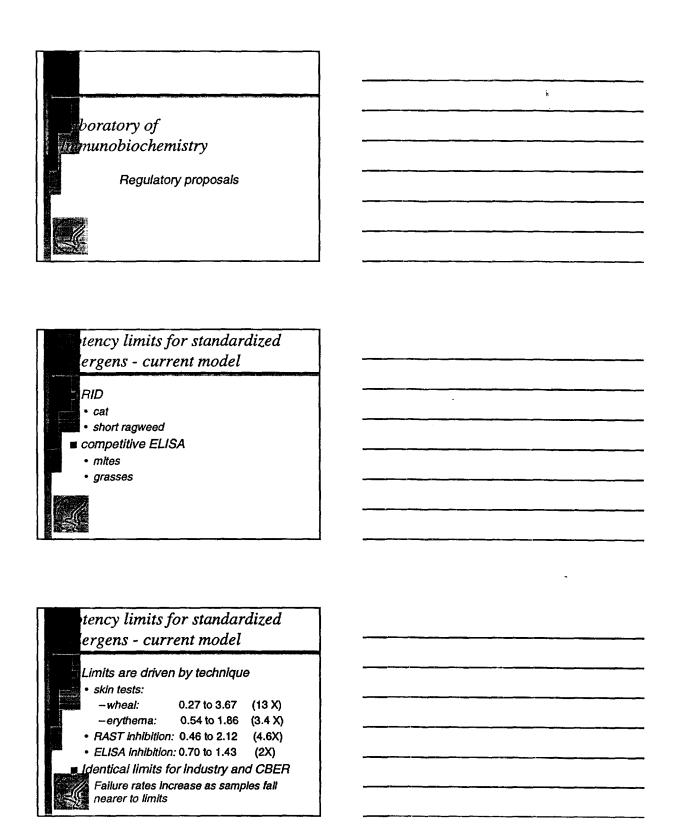


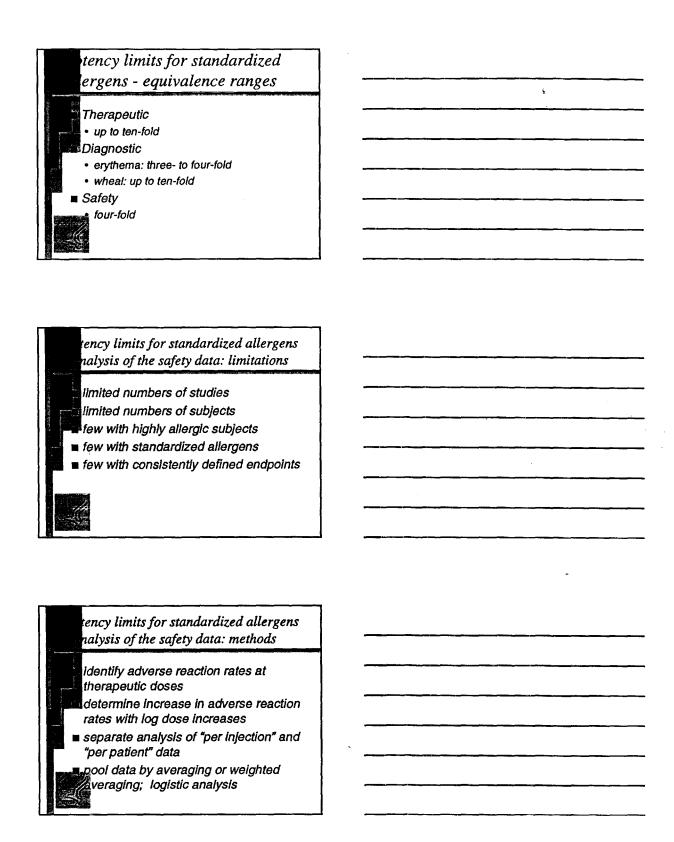


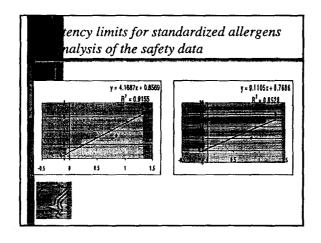


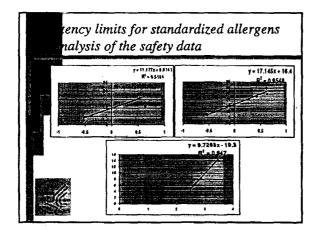
nunomodulation: lipopolysaccharides onclusions
LPS co-administered with Hev b 5/MBP accentuates • anti-Hev b 5 IgE and IgG responses • anti-Hev b 5 and anti-MBP splenocyte proliferation

nunomodulation: l irther questions	ipopolysaccharides]		
Does the effect of	functional correlate?			
powder significant				
■ Are these effects s specific?	train- or antigen-		····	
is the amount of Li				
xtracts significant	ţ			
B research pro	gram summary			
Allergen structure	■ Glycosylation		- Tent	
ano lunction	Enzyme activityIdentification methods			
■ Immunomodulation	■ Epitopes■ DNA vaccines			
	■ LPS ■ Cross-sensitization			

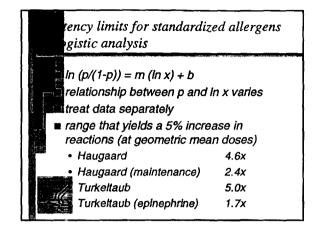








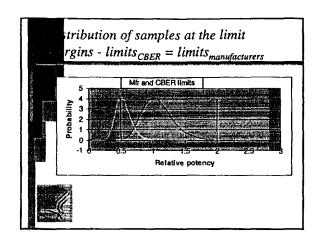
	tency limits for . nalysis of the sa	standardized allergens fety data
	saysis of the su	,
	Averages	siona (A percent/A log dosa ± SE)
	 per patient: 	13.4 ± 3.7
	• per injection:	8.2 ± 2.1
	• all data:	10.3 ± 2.1
	Weighted avera	ges
	 per patient: 	13.6 ± 5.5
March Company	per injection:	5.9 ± 4.3
16	all data:	9.3 ± 3.4
S		

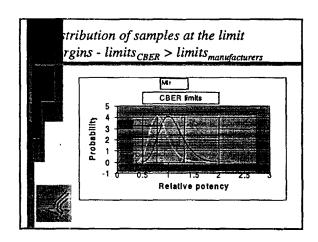


what is the α of the products that are sent to us? How does it compare to the assay? Assuming Gaussian distributions: σ(obs)² = (CBER)² + σ(manu)²

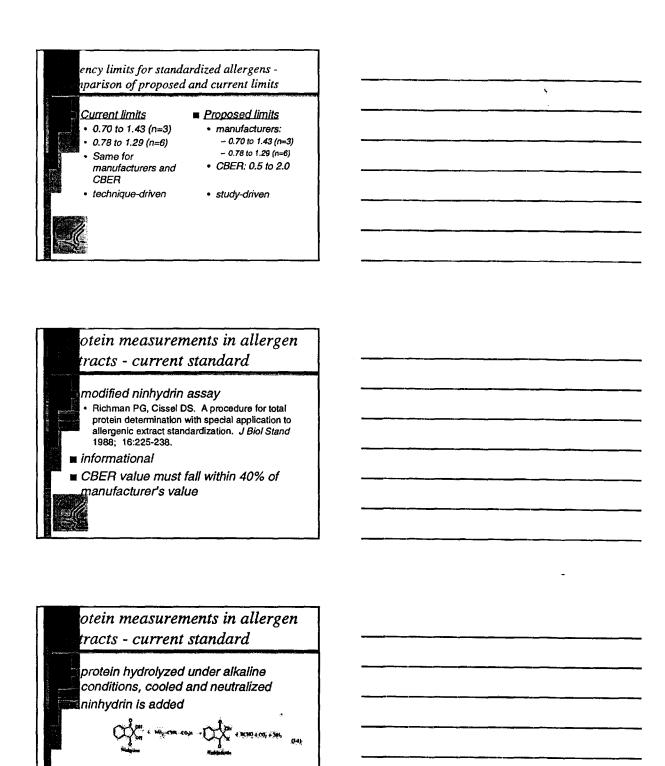
timate of σ of submitted products From 1995-1997, 53/414 or 13% of extracts falled. $\sigma(obs) = 0.12$ $\sigma(CBER) = 0.1375/\sqrt{3} = 0.08$ $\sigma(manu) = \sqrt{\sigma(obs)^2} - \sigma(CBER)^2$ $\sigma(manu) = 0.092$

w tightly should we regulate ergens? If the σ of the products that are sent to us is high, we need to insist on equivalence to reference at α On the other hand, if the σ of the products that are sent to us is low, we need to test at poundaries to eliminate utliers tency limits for standardized allergens kelihood of lot differences For Gaussian **■** When $\sigma = 0.1$ • r_{mean} = 0.798 * 0.1 distribution • r_{mean} = 0.798σ = 0.08 [log] · r_{95%} = 2.77 σ = 1.2 $r_{\it gSX} = 2.77 * 0.1$ = 0.28 [log] tency limits for standardized ergens - new limit proposal CBER limits: 0.5 to 2.0 Manufacturer internal limits: unchanged • for N = 3, 0.7 to 1.43 • for N = 6, 0.78 to 1.29

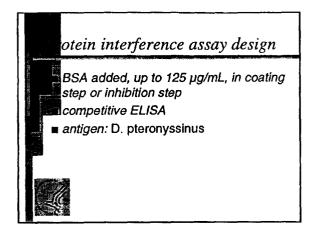


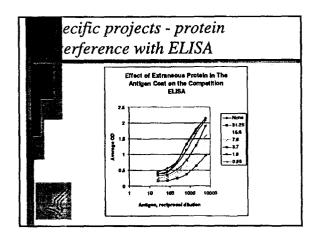


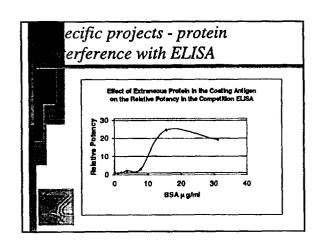
ke	elihood d	of produc	t failure	
	N(manu)	= 3	N(manu)	= 6
	RP	P(pass)	RP	P(pass)
	0.5	0.500	0.5	0.500
	0.6	0.760	0.6	0.792
	0.699	0.902	0.7	0.934
	0.7	0.903	0.776	
	0.8	0.965	8.0	0.982
	1	0.993	1	0.998
	1.2	0.976	1.2	0.989
	1.3	0.952	1.288	0.975
l	1.4	0.916	1.3	0.973
	1.431	0.902	1.4	0.944
	1.6	0.806	1.6	0.841
1/2	1.8	0.658	1.8	0.681
2/1/	2	0.500	1 2	0.500

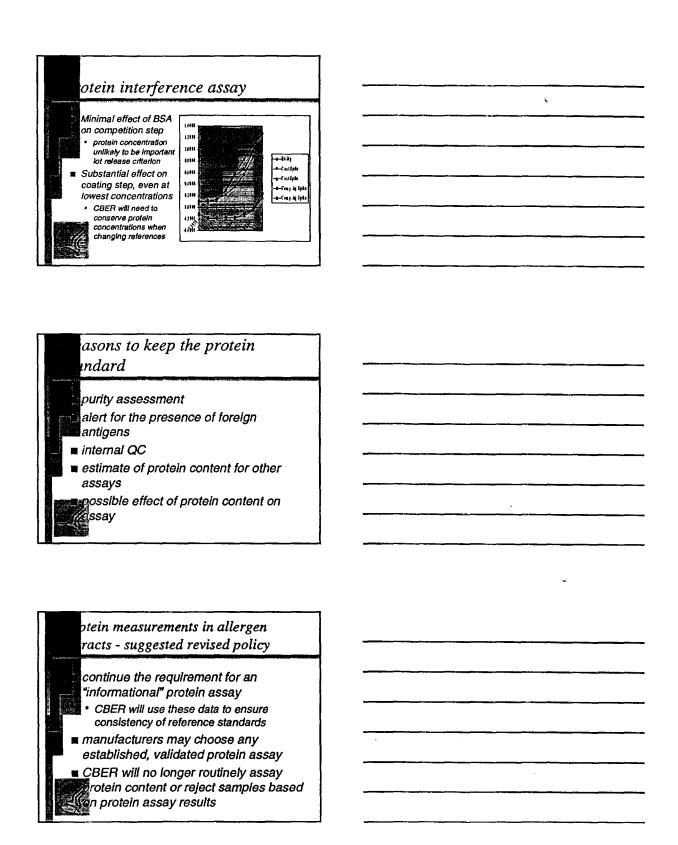


asons to keep a protein andard purity assessment alert for the presence of foreign antigens internal QC estimate of protein content for other assays cossible effect of protein content on ssay	
otein measurements in allergen racts - problems with ninhydrin	
cumbersome assay may be overly sensitive	
	-
otein measurements in allergen racts - problems with other methods	
glycerol may affect assay	
other chemicals may affect assay	
(tyrosine, tryptophan and cysteine)	









otein measurements in allergen tracts - revision Advantages ■ Disadvantages data will be, within a ■ protein data will not be given manufacturer, comparable among the internally comparable different manufacturers ■ LIB will not replicate data as part of routine lot release no possibility of lot failure based on the otein standard - clarification This recommendation applies to standardized mite and grass allergen vaccines only. Standardized hymenoptera venoms will continue to be assayed by the ninhydrin assay as currently required. ■ The results of protein assays performed on standardized mite and grass allergen vaccines may not be used in product labeling naterials. B objectives 1999-2000 Regulatory activities ■ Research activities • glycoproteins continued staff stability/expansion acid phosphatase active improvement of MALDI-TOF support program for standardized allergens • Hev b 5 epitopes DNA vaccines support for future lipopolysaccharides standardization efforts · latex cross sensitization (per Advisory Committee recommendations)

Abbreviations:

αα

amino acid residues

APMA

Allergen Product Manufacturers Association

BALB/c

an inbred strain of albino mice

BSA

bovine serum albumin

Bv

baculovirus

CBER

the Center for Biologics Evaluation and Research, FDA

Der f 1 and 2

allergens from the dust mite Dermatophagoides farinae

Der p 1, 2, and 5

allergens from the dust mite Dermatophagoides pteronyssinus

E5-Df

reference extract E5 from Dermatophagoides farinae

E5-Dp

reference extract E5 from Dermatophagoides pteronyssinus

E8-latex

reference extract E8 from Hevea brasiliensis latex

Ec

E. coli

ELISA

enzyme liked immunosorbent assay

Hev b 1 through 7

allergens from Hevea brasiliensis latex

Hya

hyaluronidase

IgG and IgE

immunoglobulins G and E

IT

immunotherapy

LIB

Laboratory of Immunobiochemistry, Division of Allergenic

Products and Parasitology, CBER, FDA

LPS

lipopolysaccharide

MALDI-TOF

matrix-assisted laser desorption/ionization time-of-flight mass

spectrometry

MBP

maltose binding protein

n (prefix)

native, or natural

PBS

phosphate buffered saline

RT-PCR

reverse transcriptase polymerase chain reaction

Prs a 1

an allergen from avocado (Persea americana)

QC

quality control

r (prefix)

recombinant

r mean and r 95%

the ratio of the RPs of two sequential lots of allergen extracts, from a population of extracts whose RPs are represented by a Gaussian distribution around a mean RP of 1. r_{mean} is the average of all r's; and $r_{95\%}$ is the r value below which 95% of sequential

lots will fall.

RAST

radioallergosorbent test

RID

radial immunodiffusion

RP

relative potency

 σ (obs), σ (CBER), and σ (manu)

standard deviations of the observed allergen extracts, of the CBER

assay, and of the manufacturers' submitted products

SD

standard deviation

SDS-PAGE

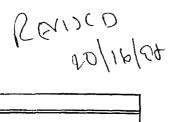
sodium dodecyl sulfate polyacrylamide gel electrophoresis

Th1

a subset of helper T-cells

TMB

colorimetric substrate for horseradish peroxidase



Reference Replacement: Track 1

E6-Dp			
	Date	# Days to Complete	Procedure
Start Date	11/1/98	7	Select Candidate(s)
	11/08/98	28	Initial CBER Testing
	12/06/98	7	Data Analysis/Select Candidate for Mfr Testing
	12/13/98	14	Hold & Purchase/Call & Send to Mfrs
	12/27/98	84	Mfr testing
	03/21/99	14	Data Analysis
	04/04/99	14	Final Decision; Memo & Shipment to Mfr
Completion Date	04/18/99	168	

E4-Or			
	Date	# Days to Complete	Procedure
Start Date	4/18/99	7	Select Candidate(s)
	04/25/99	28	Initial CBER Testing
	05/23/99	7	Data Analysis/Select Candidate for Mfr Testing
	05/30/99	14	Hold & Purchase/Call & Send to Mfrs
	06/13/99	84	Mfr testing
	09/05/99	14	Data Analysis
	09/19/99	14	Final Decision; Memo & Shipment to Mfr
Completion Date	10/03/99	168	

E4-Sv			
	Date	# Days to Complete	Procedure
Start Date	10/3/99	7	Select Candidate(s)
	10/10/99	28	Initial CBER Testing
	11/07/99	7	Data Analysis/Select Candidate for Mfr Testing
	11/14/99	14	Hold & Purchase/Call & Send to Mfrs
	11/28/99	84	Mfr testing
	02/20/00	14	Data Analysis
	03/05/00	14	Final Decision; Memo & Shipment to Mfr
Completion Date	03/19/00	168	

E4-Mf			
	Date	# Days to Complete	Procedure
Start Date	3/19/00	7	Select Candidate(s)
	03/26/00	28	Initial CBER Testing
	04/23/00	7	Data Analysis/Select Candidate for Mfr Testing
	04/30/00	14	Hold & Purchase/Call & Send to Mfrs
	05/14/00	84	Mfr testing
	08/06/00	14	Data Analysis
	08/20/00	14	Final Decision; Memo & Shipment to Mfr
Completion Date	09/03/00	168	

Reference Replacement: Track 1, Page 2

E5-Ber			
	Date	# Days to Complete	Procedure
Start Date	9/3/00	7	Select Candidate(s)
	09/10/00	28	Initial CBER Testing
	10/08/00	7	Data Analysis/Select Candidate for Mfr Testing
	10/15/00	14	Hold & Purchase/Call & Send to Mfrs
	10/29/00	84	Mfr testing
	01/21/01	14	Data Analysis
	02/04/01	14	Final Decision; Memo & Shipment to Mfr
Completion Date	02/18/01	168	

E7-Df			
	Date	# Days to Complete	Procedure
Start Date	2/18/01	7	Select Candidate(s)
	02/25/01	28	Initial CBER Testing
	03/25/01	7	Data Analysis/Select Candidate for Mfr Testing
	04/01/01		Hold & Purchase/Call & Send to Mfrs
	04/15/01	84	Mfr testing
	07/08/01	14	Data Analysis
	07/22/01	14	Final Decision; Memo & Shipment to Mfr
Completion Date	08/05/01	168	

Reference Replacement: Track 2

C7-Cat			
	Date	# Days to Complete	Procedure
Start Date	11/1/98	7	Select Candidate
	11/08/98	14	Initial CBER Testing
	11/22/98	28	Dilute for Std Curve & Test
	12/20/98	7	Data Analysis
	12/27/98	7	Call & Send to Mfrs
	01/03/99	84	Mfr testing
	03/28/99	14	Data Analysis
	04/11/99	14	Final Decision; Memo & Shipment to Mfr
Completion Date	04/25/99	175	

E4-Rt			
	Date	# Days to Complete	Procedure
Start Date	4/25/99	7	Select Candidate(s)
	05/02/99	28	Initial CBER Testing
	05/30/99	7	Data Analysis/Select Candidate for Mfr Testing
	06/06/99	14	Hold & Purchase/Call & Send to Mfrs
	06/20/99	84	Mfr testing
	09/12/99	14	Data Analysis
	09/26/99	14	Final Decision; Memo & Shipment to Mfr
Completion Date	10/10/99	168	

C11-Ras			
	Date	# Days to Complete	Procedure
Start Date	10/10/99	7	Select Candidate
	10/17/99	14	Initial CBER Testing
	10/31/99	28	Dilute for Std Curve & Test
	11/28/99	7	Data Analysis
	12/05/99	7	Call & Send to Mfrs
	12/12/99	84	Mfr testing
•	03/05/00	14	Data Analysis
	03/19/00	14	Final Decision; Memo & Shipment to Mfr
Completion Date	04/02/00	175	

E7-Ti			
	Date	# Days to Complete	Procedure
Start Date	4/2/00	7	Select Candidate(s)
	04/09/00	28	Initial CBER Testing
	05/07/00	7	Data Analysis/Select Candidate for Mfr Testing
	05/14/00	14	Hold & Purchase/Call & Send to Mfrs
	05/28/00	84	Mfr testing
	08/20/00	14	Data Analysis
	09/03/00	14	Final Decision; Memo & Shipment to Mfr
Completion Date	09/17/00	168	

Reference Replacement: Track 2, Page 2

E12-Rye			
	Date	# Days to Complete	Procedure
Start Date	9/17/00	7	Select Candidate(s)
	09/24/00	28	Initial CBER Testing
	10/22/00	7	Data Analysis/Select Candidate for Mfr Testing
	10/29/00	14	Hold & Purchase/Call & Send to Mfrs
	11/12/00	84	Mfr testing
	02/04/01	14	Data Analysis
	02/18/01	14	Final Decision; Memo & Shipment to Mfr
Completion Date	03/04/01	168	

E5-Jkb			
	Date	# Days to Complete	Procedure
Start Date	3/4/01	7	Select Candidate(s)
	03/11/01	28	Initial CBER Testing
	04/08/01	7	Data Analysis/Select Candidate for Mfr Testing
	04/15/01	14	Hold & Purchase/Call & Send to Mfrs
	04/29/01	84	Mfr testing
	07/22/01	14	Data Analysis
	08/05/01	14	Final Decision; Memo & Shipment to Mfr
Completion Date	08/19/01	168	

Release limits data - log 10 analyses

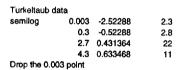
Therapeutic range data only

Haugaard data total data semilog			
	0.7	-0.1549	0.56
	7	0.845098	3.3
	21	1.322219	7.1

m	b	4.168735	0.856925
Sem	SEb	1.266292	1.152824
r2	SEv	0.915525	1.349959
F	df	10.8378	1
ssreg	sesresid	19.75068	1.822389

Haugaard data maintenance only semilog

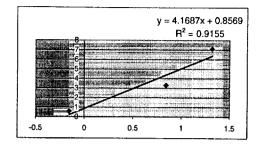
	0.7	-0.1549	0.4
	7	0.845098	5.24
	21	1.322219	19
m	b	9.110539	0.768604
Sem	SEb	3.784991	3.445832
r2	SEy	0.852805	4.035075
F	df	5.793732	1
ssreg	sesresid	94.33257	16.28183

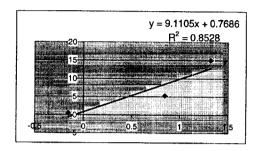


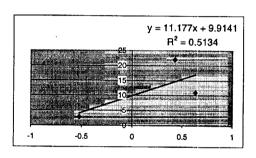
m	b	11.17747	9.914111
Sem	SEb	10.88098	5.828387
r2	SEy	0.513438	9.503621
F	df	1.055238	1
ssreg	sesresid	95.30786	90.31881

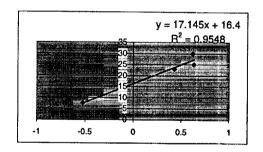
Turkeltaub epi data semilog 0.3 -0.52288 7.5 0.82 -0.08619 15 2.7 0.431364 23 4.2 0.623249 4.3 0.633468 25

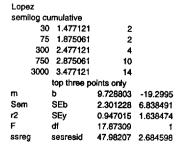
m	b	17.14537	16.39997
Sem	SEb	2.153875	1.079789
r2	SEy	0.954796	2.179326
F	df	63.36544	3
ssreg	sesresid	300.9516	14.24838

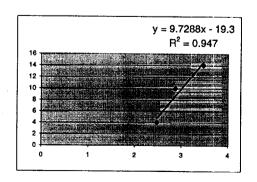












Equal weight averages

Per injection summary

i di injediton danimarj		
	slope	SE
Haugaard all data	4.17	1.27
Haugaard(all data	9.11	3.78
Turkeltaub worst case	11.18	10.88
Per patient summary		
	slope	SE
Turkeltaub epi data	17.15	2.15
Lopez worst case	9.73	2.30

All data

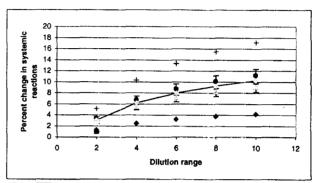
2	4	6	8	10
1.254914	2.509828	3.243906	3.764743	4.168735
2.742546	5.485091	7.089377	8.227637	9.110539
3.364753	6.729507	8.697761	10.09426	11.17747
5.16127	10.32254	13.34169	15.48381	17.14537
2.928662	5.857323	7.57048	8.785985	9.728803
3.090429	6.180858	7.988643	9.271287	10.26618
1.403562	2.807124	3.628155	4.210685	4.662531
0.627692	1.255384	1.62256	1.883076	2.085147
	1.254914 2.742546 3.364753 5.16127 2.928662 3.090429 1.403562	1.254914 2.509828 2.742546 5.485091 3.364753 6.729507 5.16127 10.32254 2.928662 5.857323 3.090429 6.180858 1.403562 2.807124	1.254914 2.509828 3.243906 2.742546 5.485091 7.089377 3.364753 6.729507 8.697761 5.16127 10.32254 13.34169 2.928662 5.857323 7.57048 3.090429 6.180858 7.988643 1.403562 2807124 3.628155	2 4 6 8 1.254914 2.509828 3.243906 3.764743 2.742546 5.485091 7.089377 8.227637 3.364753 6.729507 8.697761 10.09426 5.16127 10.32254 13.34169 15.48381 2.928662 5.857323 7.57048 8.785985 3.090429 6.180858 7.989863 9.271287 1.403562 2.807124 3.628155 4.210685 0.627692 1.255384 1.62256 1.883076

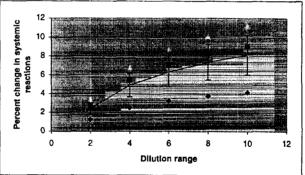
per injection only

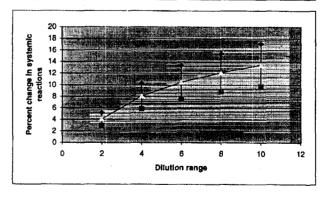
	2	4	6	8	10
	1.254914	2.509828	3.243906	3.764743	4.168735
	2.742546	5.485091	7.089377	8.227637	9.110539
	3.364753	6.729507	8.697761	10.09426	11.17747
Average	2.454071	4.908142	6.343682	7.362213	8.152248
SD	1.084098	2.168196	2.802353	3.252294	3.601296
SE	0.625904	1.251809	1.617939	1.877713	2.079209

per patient only M

	2	4	6	8	10
	5.16127	10.32254	13.34169	15.48381	17.14537
	2.928662	5.857323	7.57048	8.785985	9.728803
Average	4.044966	8.089932	10.45609	12.1349	13.43709
SD	1.578693	3.157385	4.080861	4.736078	5.244303
SE	1.116304	2.232609	2.885605	3.348913	3,708283







Weighted averages

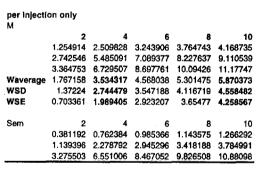
Per injection summary		
	slope	SE
Haugaard all data	4.17	1.27
Haugaard(all data	9.11	3.78
Turkeltaub worst case	11.18	10.88

Per patient summary

		slope	SE
Turkelta	ub epi data	17.15	2.15
Lopez	worst case	9.73	2.30

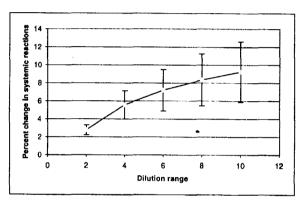
All data

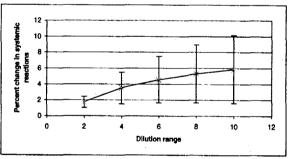
М	2	4	6	8	10
	1.254914	2.509828	3.243906	3.764743	4.168735
	2.742546	5.485091	7.089377	8.227637	9.110539
	3.364753	6.729507	8.697761	10.09426	11.17747
	5.16127	10.32254	13.34169	15.48381	17.14537
	2.928662	5.857323	7.57048	8.785985	9.728803
Waverage	2.784719	5.569439	7.198395	8.354158	9.250637
WSD	1.444579	2.889158	3.734183	4.333737	4.798788
WSE	0.554291	1.567771	2.303663	2.880179	3.356007
Sem	2	4	6	8	10
	0.381192	0.762384	0.985366	1.143575	1.266292
	1.139396	2.278792	2.945296	3.418188	3.784991
	3.275503	6.551006	8.467052	9.826508	10.88098
	0.648381	1.296762	1.676041	1.945143	2.153875
	0.692739	1.385477	1.790703	2.078216	2.301228

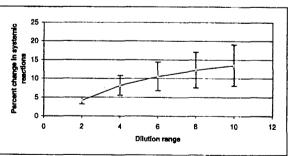


per patient only M

	2	4	6	8	10
	5.16127	10.32254	13.34169	15.48381	17.14537
	2.928662	5.857323	7.57048	8.785985	9.728803
Waverage	4.081888	8.163775	10.55153	12.24566	13.55974
WSD	1.579556	3.159112	4.083093	4.738668	5.247171
WSE	0.914116	2.585509	3.799115	4.749884	5.534603
Sem	2	4	6	8	10
	0.648381	1.296762	1.676041	1.945143	2.153875
	0.692739	1.385477	1.790703	2.078216	2.301228





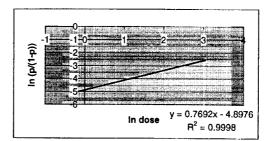


Logistic analysis

Haugaard data
total data

dose	In dose	p (%)	р	In(p/(1-p))
0.7	-0.356675	0.56	0.0056	-5.179373
7	1.94591	3.3	0.033	-3.377691
21	3.044522	7.1	0.071	-2.571429

m	b	0.769178	-4.897559
Sem	SEb	0.011772	0.024677
r2	SEy	0.999766	0.028897
F	df	4269.166	1
ssreg	sesresid	3.564995	0.000835



Haugaard data maintenance only

m Sem r2 F

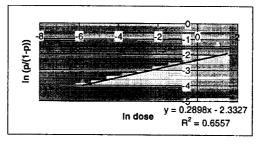
ssreg

d	ose	in dose	p (%)	р	In(p/(1-p))
	0.7	-0.356675	0.4	0.004	-5.517453
	7	1.94591	5.24	0.0524	-2.895026
	21	3.044522	15	0.15	-1.734601
b		1.116305	-5.10659		
SEb		0.020031	0.041989		
SEy		0.999678	0.049169		
df		3105.859	1		
sesresid		7.508808	0.002418		

((d-1)/d) u
a) 4 4 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
- <u>- 5 2 </u>

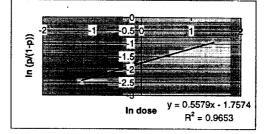
Turkeltaub	dose	In dose	p (%)	р	In(p/(1-p))
	0.003	-5.809143	2.3	0.023	-3.748992
	0.3	-1.203973	2.8	0.028	-3.547151
	2.7	0.993252	22	0.22	-1.265666
	4.3	1.458615	11	0.11	-2.090741

m	b	0.289809	-2.332665
Sem	SEb	0.148506	0.459589
r2	SEy	0.655667	0.854506
F	df	3.80833	2
ssreg	sesresid	2.780765	1.460359



Turkeltaub epi data

dose	In dose	p (%)	р	ln(p/(1-p))
0.3	-1.203973	7.5	0.075	-2.512306
0.82	-0.198451	15	0.15	-1.734601
2.7	0.993252	23	0.23	-1.208311
4.2	1.435085	30	0.3	-0.847298
4.3	1.458615	25	0.25	-1.098612



m	b ·	0.557871	-1.757435
Sem	SEb	0.061092	0.070521
r2	SEy	0.965272	0.142332
F	df	83.38634	3
ssreg	sesresid	1.689284	0.060776

$\ln (p/(1-p)) = m(\ln x) + b$

Study Haugaard	geo mean	m	b	In p/(1-p)	p/(1-p)	р	p + .05	In p/(1-p)	new dose	factor
	4.69	0.77	-4.90	-3.71	0.02	0.02	0.07	-2.53	21.77	4.64
Haugaard (n	naintenance)									
	4.69	1.12	-5.11	-3.38	0.03	0.03	0.08	-2.40	11.26	2.40
Turkeltaub										
	1.52	0.29	-2.33	-2.21	0.11	0.10	0.15	-1.75	7.59	5.01
Turkeltaub (e	epi)									
	1.64	0.56	-1.76	-1.48	0.23	0.19	0.24	-1.18	2.82	1.72



DEPARTMENT OF HEALTH & HUMAN SERVICES FDA/CBER/OVRR/DAPP

Memorandum

Date

DRAFT

To

From

Jay E. Slater, MD, Chief, Laboratory of Immunobiochemistry

Through

Subject

Elimination of the requirement for the ninhydrin total protein assay for

standardized mite and grass allergen vaccines

The determination of the protein content of allergen extracts has been used as a lot release criterion for standardized allergen vaccines. With advances in allergen standardization and identification, the protein content of an individual vaccine may be of questionable relevance to the safety and efficacy of the product. The purpose of this memorandum is to discuss the advantages and disadvantages of changing the requirement that manufacturers determine the protein content of standardized mite and grass allergen vaccines, and to recommend a specific diminution of the regulatory requirements for this assay.

Reasons for a protein lot release requirement

The most important reason for determining total protein is as a measure of product consistency from lot to lot. Large variations in protein content may signal manufacturing deficiencies that warrant attention. Furthermore, a decrease in the potency/unit protein of an allergen extract may be a sensitive indicator of allergen degradation or contamination.

Another reason to monitor the protein content is the need to estimate the amount of material needed for other laboratory assays of possible regulatory interest. These include analysis by gel electrophoresis, immunoblot, radial immunodiffusion, isoelectric focussing, crossed radial immunoelectrophoresis, HPLC and MALDI-TOF.

Finally, contaminating proteins may interfere with other assays. Primarily, these may affect protein-protein interactions, especially the solid-phase coating step in ELISA-based assays. In addition, other reactions depending on antigen-antibody interactions may be susceptible to unanticipated effects contributed by extraneous proteins.

The choice of a standard protein assay for allegen extracts

Unfortunately, each of the common protein assays has limitations that are of special concern when evaluating allergen vaccines. Glycerol, a common component of allergen preparations, interferes with Lowry-based assays (including the BCA assay), and may affect the Coomassle blue-based assays (e.g. Bradford) as well. In addition, these assays are all dependent upon the presence of particular amino acid residues for color development, which may not be present in comparable amounts in all allergens. The latex allergen Hev b 5, for instance, is devoid of tyrosine, tryptophan and cysteine¹, and may be undetectable using these techniques².

In consideration of these concerns, CBER developed and adopted a modification of the more cumbersome ninhydrin technique of protein determination 3 . In this assay, the protein is hydrolyzed under alkaline conditions. The mixture is cooled and neutralized, and ninhydrin is added. Ninhydrin elicits the oxidative deamination of the α -amino group, and it is the reduced form of ninhydrin that absorbs light near 570 nm. Unlike other methods, the reaction of ninhydrin with amino acids is largely independent of the side chain; thus, in principle, the ninhydrin assay can closely approximate the results of an amino acid analysis 4 .

It is notable that release limits were not established for the protein content of standardized allergen vaccines. Rather, the results of the ninhydrin assay have been required for information only. When the results have been checked as part of the lot release program of LIB, the requirement has been that the results of the CBER assay be within 40% of the manufacturer's result.

Problems associated with the ninhydrin assay

Compared with other available protein assays, the ninhydrin assay is lengthy and difficult. Toxic, caustic reagents are used at high temperatures. Hydrolysis is achieved by the addition of 10 N NaOH and incubation in a 150°C oven. Continued incubation of open tubes at 110°C is necessary to eliminate free ammonia and decrease background signal in the assay. The mixture is neutralized with 10 N acetic acid before the ninhydrin reagent is added³.

Furthermore, the sensitivity of the ninhydrin assay may be a limitation. While the standard protein assays are unlikely to detect small peptides and amino acids, the ninhydrin assay will do so. These small peptide sequences may be of less concern than larger proteins that are more likely to be allergenic. Furthermore, by increasing the detection of small peptides and amino acids, the ninhydrin assay may be less likely than other protein assays to detect shifts in the concentration of proteins of greater immunologic significance⁵.

Is the ninhydrin assay necessary to measure extraneous protein?

Theoretically, it is possible that an allergen submitted for lot release will be contaminated with other allergens or extraneous proteins that will be detected only by an increase in the total protein content. However, it is likely that these extraneous proteins will be detected in the identity testing. In addition, following the lot-to-lot total protein content of a product over time can be accomplished as well using one of the standard protein assays. Although glycerol may interfere with the assays, the glycerol content should be relatively stable from lot to lot.

Are total protein assays necessary to establish the amount of allergen to be used for other assays?

Knowing the amount of protein in a sample can save time and expense in running certain assays. Examples include HPLC, IEF and immunoblots, where the reagent, labor and equipment time expenditures are high. However, for ELISA assays, costs are relatively low, multiple dilutions are performed as a matter of course, and the initial concentration is less important. Furthermore, even for those assays for which the initial protein concentration would be useful information, alternative protein assays, when properly validated, can provide adequate information.

Are protein assays necessary to prevent interference with other assays?

We examined the possibility that protein levels may affect the results of the competitive ELISA. We found added protein significantly inhibited the binding of allergen to microtiter wells, but had no measurable effect on the competition step. This suggests that the protein content of an allergen vaccine submitted for lot release will not affect the results of lot release testing. However, the protein content of the candidate allergen vaccine must be considered when CBER selects a reference extract (which is, in all cases, the allergen bound to the wells in the initial step). Once again, alternative protein assays, when properly validated, can be used to monitor the protein content for this purpose.

Thus the ninhydrin assay currently required by CBER for the approval of standardized allergen vaccines is difficult to use, and the assay sensitivity may be of limited utility or relevance. It is probably no longer necessary as a quality control measure, and other protein assays are probably sufficient indicators of extraneous protein content. Initial estimates of protein content are not needed for the radial immunodiffusion or competitive ELISA assays, and the estimates provided by the standard protein assays would be sufficiently accurate for the establishment of initial conditions for tests such as HPLC and IEF. Finally, the protein content of an extract

submitted for lot release does not appear to affect the results of the competitive ELISA assay used by CBER.

Current options (summarized in Table)

- 1. No change. This is the most conservative approach. The main advantage is that CBER will continue to require that each manufacturer utilize the same assay method, and that information on the specific activity (relative potency/unit protein) of allergen vaccines from different manufacturers will be directly comparable. Another advantage is that CBER biologists will need to master only one, albeit difficult, protein assay. CBER has extensive experience with this assay, and has collected large amounts of data from the manufacturers since the initiation of the standardization program. These data could be utilized to initiate future studies of the potency of allergen vaccines. The disadvantage of this approach is that CBER will continue to require the manufacturers to perform a difficult assay using hazardous reagents to collect data that are, at best, of uncertain value. We have no example yet of a production or manufacturing defect that has been uncovered as a result of the data obtained using the protein assay.
- 2. Require the ninhydrin assay, but eliminate limits on the protein content. At present, the only limits that are in force reflect on the accuracy of the assay and of the laboratory personnel who perform the assay, in the manufacturers' laboratories and at CBER. The advantage of this approach is that we will continue to collect comparable, reliable data, which can be used for quality control and assay dosing purposes. However, we eliminate the possibility of lot failure based on the assay, and eliminate the need for LIB to perform the assay as part of the lot release process.
- 3. Require a protein assay, permit the choice of any standard protein assay, but set limits on the protein content. Under this option, we expect that, once an assay is chosen, it will be changed only with adequate justification. The main advantage of this approach is that we will continue to check on the protein content of the vaccines and the accuracy of the manufacturers' laboratory determinations. We will also continue to collect data on allergen vaccines that are, within a given manufacturer, internally comparable. In addition, these data will be helpful in determining initial amounts of allergen to use in various subsequent assays. There is at least a theoretical possibility that the standard protein assays measure larger, more significant allergenic proteins, while the ninhydrin assay measures all proteins and polypeptides. The major disadvantages are that LIB will have to run each of several different protein assays on a routine basis, and the acceptable variations will have to be determined individually for each assay. The protein data will not be comparable among the different manufacturers. Lot release failure on the basis of the protein assay remains a possibility in this option, and it is not clear that such failure has any justifiable basis.
- 4. Require any protein assay, but eliminate limits on the protein content. The advantage of this approach is that we continue to collect data on allergen vaccines that are, within a given manufacturer, internally comparable. These data will be helpful in determining initial amounts of allergen to use in various assays. LIB personnel will only need to replicate the manufacturer's data as part of the investigation of a specific problem, but not as part of routine lot release. Once again, we eliminate the possibility of lot failure based on the assay. However, the protein data will not be comparable among the different manufacturers.
- 5. Eliminate protein assay requirement. The advantage of this approach is that it relieves the industry and LIB of all regulatory burdens associated with the protein content of allergen vaccines. There are several disadvantages. The major disadvantage is that CBER will no longer have any information on the protein content of standardized allergen vaccines, unless the manufacturers voluntarily provide this information. Of lesser importance is that LIB workers will no longer have an initial estimate from the manufacturer of protein content for assays such as IEF, HPLC and MALDI-TOF, and for the selection of appropriately consistent reference standards.

Recommendation

Overall, there appears to be little justification for continuing to require the use of the ninhydrin assay. However, manufacturers should continue to perform protein assays on each lot of material, and CBER should require this information as part of its lot release program. The choice of a protein assay will be left to the manufacturer, which must provide proficiency and validation data on the particular assay, and an SOP that can be followed by CBER personnel should replication be required. Since one of the reasons to require protein data is for ongoing monitoring of particular allergen vaccines, manufacturers will be expected to use one protein assay consistently unless compelling reasons are presented to CBER.

Likewise, the advantages of continuing to require that CBER personnel replicate the manufacturer's data within 40% limits are uncertain. This does not represent a true limit on protein content, but rather a test of the accuracy of the manufacturer's assay technique. Appropriate proficiency and validation should be adequate for this purpose.

At this time, there are no data to support the establishment of a true limit on the protein content of allergen vaccines.

The major substantive disadvantage of this option (#4, above) is that the protein data will not be comparable among the different manufacturers. We have no evidence that such data have been used in the past. Furthermore, if a particular issue should arise for which these data are needed, CBER personnel can determine the protein content of allergen vaccines from different manufacturers by performing one of the protein assays (including, at the Lab Chief's discretion, the ninhydrin assay) on lot release samples.

This recommendation applies to standardized mite and grass allergen vaccines only. Standardized hymenoptera venoms will continue to be assayed by the ninhydrin assay as currently required. Furthermore, the results of protein assays performed on standardized mite and grass allergen vaccines may not be used in product labeling materials.

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Approach	Lot-to-lot consistency	Industry-wide data	Initial quantities for	Control for possible	Assay is easy to perform		Measures mostly
			assays	interference with other assays	Mfr	LIB	proteins >10 kDa
1. Status quo	Yes	Yes	Yes	Yes	No	No	No
2. Ninhydrin/no limits	Yes	Yes	Yes	Yes	No	N/A	No
3. Any assay/limits	Yes	No	Yes	Yes	Yes	Yes	Yes
4. Any assay/no limits	Yes	No	Yes	Yes	Yes	N/A	Yes
5. No requirement	No	No	No	No	N/A	N/A	No

^{*} However, LIB staff will have to perform multiple protein assays, and variance limits will have to be set for each assay.